

SEQUENCE LISTING

<110> Griffith, Than J et al.

<120> T CELL EPITOPES OF RYEGRASS POLLEN ALLERGEN

<130> IMI-040CP3

<140> 08/737,904

<141> 1996-11-20

<150> 08/106,016

<151> 1993-08-13

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<170> PatentIn Ver. 2.0

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(212> DNA

<213> Escherichia coli

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⊉21> CDS

(222> (40)..(942)

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tac acg gtg gct cta ttc ctc gcc gtg gcc ctc gtg gcg ggc ccg gcc 102

Tyr Thr Val Ala Leu Phe Leu Ala Val Ala Leu Val Ala Gly Pro Ala

10 20

gcc tcc tac gcc gct gac gcc ggc tac adc ccc gca gcc gcg gcc acc 150
Ala Ser Tyr Ala Ala Asp Ala Gly Tyr Thr Pro Ala Ala Ala Ala Thr
25 30 35

ccg gct act cct gct gcc acc ccg gct gcg gct gga ggg aag gcg acg 198
Pro Ala Thr Pro Ala Ala Thr Pro Ala Ala Gly Gly Lys Ala Thr
40
45
50

acc gac gag cag aag ctg ctg gag gac gtc aac gct ggc ttc aag gca 246
Thr Asp Glu Gln Lys Leu Leu Glu Asp Val Asn Ala Gly Phe Lys Ala 55 60 65

gcc gtg gcc gcc gct gcc aac gcc cct ccg gcg gac aag ttc aag atc 294
Ala Val Ala Ala Ala Ala Asn Ala Pro Pro Ala Asp Lys Phe Lys Ile
70 75 80 85

ttc gag gcc gcc ttc tcc gag tcc tcc aag ggc ctc ctc gcc acc tcc 342
Phe Glu Ala Ala Phe Ser Glu Ser Ser Lys Gly Leu Leu Ala Thr Ser
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gec gec aag gea eee gge etc ate eee aag etc gae ace gee tae gae 390

47

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					ctc Leu											486
					gtc Val 155											534
					gag Glu											582
	_			_	acc Thr	_	_		_	_				-		630
					agt Ser											678
					acc Thr											726
					gcc Ala 235											774
					gcc Ala											822
					ggc Gly											870
					ggc											918
	_	_		-	tac Tyr		-	tgat	cago	ctt q	gctaa	atata	ac ta	actga	aacgt	972
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tcatgcagcc gcgatcgaga gggcttgcat gcttgtaata attcaatatt tttcatttct										1092						
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Val Ala Gly Pro Ala Ala Ser Tyr Ala Ala Asp Ala Gly Tyr Thr Pro 20 25 30

Ala Ala Ala Ala Thr Pro Ala Thr Pro Ala Ala Thr Pro Ala Ala Ala 35 40 45

Gly Gly Lys Ala Thr Thr Asp Glu Gln Lys Leu Leu Glu Asp Val Asn 50 55 60

Ala Gly Phe Lys Ala Ala Val Ala Ala Ala Ala Asn Ala Pro Pro Ala 65 70 75 80

Asp Lys Phe Lys Ile Phe Glu Ala Ala Phe Ser Glu Ser Ser Lys Gly 85 90 95

Leu Leu Ala Thr Ser Ala Ala Lys Ala Pro Gly Leu Ile Pro Lys Leu 100 105 110

Asp Thr Ala Tyr Asp Val Ala Tyr Lys Ala Ala Glu Gly Ala Thr Pro 115 120 125

Glu Ala Lys Tyr Asp Ala Phe Val Thr Ala Leu Thr Glu Ala Leu Arg 130 135 140

Val Ile Ala Gly Ala Leu Glu Val His Ala Val Lys Pro Ala Thr Glu 145 150 155 160

Glu Val Pro Ala Ala Lys Ile Pro Thr Gly Glu Leu Gln Ile Val Asp 165 170 175

Lys Ile Asp Ala Ala Phe Lys Ile Ala Ala Thr Ala Ala Asn Ala Ala 180 185 190

Pro Thr Asn Asp Lys Phe Thr Val Phe Glu Ser Ala Phe Asn Lys Ala 195 200 205

Leu Asn Glu Cys Thr Gly Gly Ala Tyr Glu Thr Tyr Lys Phe Ile Pro 210 215 220

Ser Leu Glu Ala Ala Val Lys Gln Ala Tyr Ala Ala Thr Val Ala Ala 225 230 235 240

Ala Pro Glu Val Lys Tyr Ala Val Phe Glu Ala Ala Leu Thr Lys Ala

245 250 255

Ile Thr Ala Met Thr Gln Ala Gln Lys Ala Gly Lys Pro Ala Ala Ala 260 265 270

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Ala Gly Ala Ala Thr Ala Ala Ala Gly Gly Tyr Lys Ala 290 295 300

<210> 3

<211> 20

<212> PRT

<213> Escherichia coli

<220>

<223> all occurrences of Xaa=hydroxyproline

<400> 3

Ala Asp Ala Gly Tyr Thr Xaa Ala Ala Ala Ala Thr Xaa Ala Thr Xaa 1 5 10 15

Ala Ala Thr Xaa 20

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<220>

<223> all occurrences of Xaa=hydroxyproline

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Ala Thr Thr Asp

20

<210> 5

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<223> all occurrences of Xaa =hydroxyproline

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1 5 10 15

Asp Val Asn Ala

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Ala Ala Ala Ala
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<213> Escherichia coli
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Gly Phe Lys Ala Ala Val Ala Ala Ala Ala Asn Ala Pro Pro Ala Asp
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Glu Ser Ser Lys
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Ala Ala Glu Gly
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Tyr Asp Ala Phe
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Ala Thr Pro Glu Ala Lys Tyr Asp Ala Phe Val Thr Ala Leu Thr Glu
Ala Leu Arg Val
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                                     10
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Val His Ala Val
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20

<211> 20

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<210> 15
<211> 20
<212> PRT
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<400> 15
Ile Ala Gly Ala Leu Glu Val His Ala Val Lys Pro Ala Thr Glu Glu
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Val Pro Ala Ala
<210> 16
<211> 20
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<213> Escherichia coli
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Lys Pro Ala Thr Glu Glu Val Pro Ala Ala Lys Ile Pro Thr Gly Glu
                                     10
Leu Gln Ile Val
             20
<210> 17
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<212> PRT
<213> Escherichia coli
<400> 17
Lys Ile Pro Thr Gly Glu Leu Gln Ile Val Asp Lys Ile Asp Ala Ala
Phe Lys Ile Ala
<210> 18
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Asp Lys Ile Asp Ala Ala Phe Lys Ile Ala Ala Thr Ala Ala Asn Ala
Ala Pro Thr Asn
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Ala Thr Ala Ala Asn Ala Ala Pro Thr Asn Asp Lys Phe Thr Val Phe
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                                      10
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Asp Lys Phe Thr Val Phe Glu Ser Ala Phe Asn Lys Ala Leu Asn Glu
                                      10
Cys Thr Gly Gly
<210> 21
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Asn Lys Ala Leu Asn Glu Cys Thr Gly Gly Ala Tyr Glu Thr Tyr Lys
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Phe Ile Pro Ser
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Gln Ala Tyr Ala
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1 5 10 15

Pro Glu Val Lys 20

<210> 24

<211> 20

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Ala Leu Thr Lys 20

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Gln Ala Gln Lys 20

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<400> 26

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Ala Ala Ala Thr 20

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<400> 27

Ala Gly Lys Pro Ala Ala Ala Ala Ala Thr Gly Ala Ala Thr Val Ala 1 5 10 15

Thr Gly Ala Ala

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Gly Ala Ala Thr Val Ala Thr Gly Ala Ala Thr Ala Ala Gly Ala
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Ala Thr Ala Ala
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Ile Ala Lys Val Pro Pro Gly Pro Asn Ile Thr Ala Glu Tyr Gly Asp
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Lys Trp Leu Asp
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Lys Trp Leu Asp
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10
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Asp Lys Ala Pro
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Asp Lys Ala Pro
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Glu Glu Pro Ile
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Leu Ser Gly His
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Ala Gly Glu Leu
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<210> 44
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Pro Thr Phe His
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Val Asp Gly Asp
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Val Glu Lys Gly Ser Asn Pro Asn Tyr Leu Ala Ile Leu Val Lys Tyr
Val Asp Gly Asp
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Asp Ile Lys Glu
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Ile Glu Leu Lys
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<210> 50

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Trp Arg Ile Asp
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Gly Gly Thr Lys
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Ile Pro Glu Gly
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<223> all occurrences of Xaa=hydroxyproline

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						gcg Ala										202
aag Lys	gcg Ala	gcg Ala	acc Thr 30	gag Glu	gag Glu	cag Gln	aag Lys	ctg Leu 35	atc Ile	gag Glu	aag Lys	atc Ile	aac Asn 40	gcc Ala	ggc Gly	250
ttc Phe	aag Lys	gcc Ala 45	gcc Ala	gtg Val	gcg Ala	gcc Ala	gcc Ala 50	gcg Ala	ggc Gly	gtc Val	ccg Pro	cca Pro 55	ggc Gly	gac Asp	aag Lys	298
tac Tyr	aag Lys 60	acg Thr	ttc Phe	gtc Val	gaa Glu	acc Thr 65	ttc Phe	ggc Gly	aag Lys	gcc Ala	tcc Ser 70	aac Asn	aag Lys	gcc Ala	ttc Phe	346
						aac Asn										394
						gcc Ala										442
cag Gln	ggc Gly	gcc Ala	acc Thr 110	ccc Pro	gag Glu	gcc Ala	aag Lys	tac Tyr 115	gac Asp	gcc Ala	tac Tyr	gtc Val	gcc Ala 120	acc Thr	ctc Leu	490
agc Ser	gag Glu	gcg Ala 125	ctc Leu	cgc Arg	atc Ile	atc Ile	gcc Ala 130	ggc Gly	acc Thr	ctc Leu	gag Glu	gtc Val 135	cac His	gcc Ala	gtc Val	538
aag Lys	ccc Pro 140	gct Ala	gcc Ala	gag Glu	gag Glu	gtc Val 145	aag Lys	cct Pro	atc Ile	ccc Pro	gcc Ala 150	gga Gly	gag Glu	ctg Leu	cag Gln	586
atc Ile 155	gtc Val	gac Asp	aag Lys	att Ile	gac Asp 160	gtc Val	gcc Ala	ttc Phe	aga Arg	act Thr 165	gcc Ala	gcc Ala	acc Thr	gcc Ala	gcc Ala 170	634
aac Asn	gcc Ala	gcc Ala	ccc Pro	acc Thr 175	aac Asn	gac Asp	aag Lys	ttc Phe	acc Thr 180	gta Val	ttc Phe	gag Glu	acc Thr	acc Thr 185	ttt Phe	682
aac Asn	aag Lys	gcc Ala	atc Ile 190	aag Lys	gag Glu	agc Ser	acg Thr	ggc Gly 195	ggc Gly	acc Thr	tac Tyr	gag Glu	agc Ser 200	tac Tyr	aag Lys	730
ttc Phe	att Ile	ccc Pro	acc Thr	ctt Leu	gag Glu	gcc Ala	gcc Ala	gtt Val	aag Lys	cag Gln	gcc Ala	tac Tyr	gcc Ala	gcc Ala	acc Thr	778

205	210	2	215							
gtc gca tcc gcg ccg Val Ala Ser Ala Pro 220	gag gtc aag tad Glu Val Lys Tyr 225	gcc gtc ttt g Ala Val Phe G 230	gag acc gcg ctg { Glu Thr Ala Leu	826						
aaa aag gcg gtc acc Lys Lys Ala Val Thr 235	gcc atg tcc gag Ala Met Ser Glu 240	g gcc cag aag g n Ala Gln Lys G 245	gaa gcc aag ccc { Glu Ala Lys Pro 250	874						
gcc acc gcc acc ccg Ala Thr Ala Thr Pro 255	acc ccc acc gca Thr Pro Thr Ala	a act gcc gcg g a Thr Ala Ala <i>F</i> 260	, , , , , , , , , , , , , , , , , , , ,	922						
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gctagcaata tacacatcca tcatgcacat atagagctgt gtatgtatgt gcatgcatgc 1033										
cgtggcgccg cgcaagtttg ctcataatta attcttggtt ttcgttgctt gcatccacga 1										
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Thr Pro Ala Thr Pro	Ala Thr Pro Ala	a Ala Pro Gly A 20	Ala Ala Val Pro							
Ala Gly Lys Ala Ala 25	Thr Glu Glu Gl: 30	n Lys Leu Ile (35	Glu Lys Ile Asn 40							
Ala Gly Phe Lys Ala	Ala Val Ala Al	a Ala Ala Gly V	Val Pro Pro Gly							

Asp Lys Tyr Lys Thr Phe Val Glu Thr Phe Gly Lys Ala Ser Asn Lys

Ala Phe Leu Gly Asp Leu Pro Thr Asn Tyr Ala Asp Val Asn Ser Arg
75 80 85

Ala Gln Leu Thr Ser Lys Leu Asp Ala Ala Tyr Lys Leu Ala Tyr Asp

95

60

Ala Ala Gln Gly Ala Thr Pro Glu Ala Lys Tyr Asp Ala Tyr Val Ala 110 115 Thr Leu Ser Glu Ala Leu Arg Ile Ile Ala Gly Thr Leu Glu Val His Ala Val Lys Pro Ala Ala Glu Glu Val Lys Pro Ile Pro Ala Gly Glu 145 140 Leu Gln Ile Val Asp Lys Ile Asp Val Ala Phe Arg Thr Ala Ala Thr 160 Ala Ala Asn Ala Ala Pro Thr Asn Asp Lys Phe Thr Val Phe Glu Thr 175 Thr Phe Asn Lys Ala Ile Lys Glu Ser Thr Gly Gly Thr Tyr Glu Ser 195 190 Tyr Lys Phe Ile Pro Thr Leu Glu Ala Ala Val Lys Gln Ala Tyr Ala 205 210 Ala Thr Val Ala Ser Ala Pro Glu Val Lys Tyr Ala Val Phe Glu Thr 225 Ala Leu Lys Lys Ala Val Thr Ala Met Ser Glu Ala Gln Lys Glu Ala Lys Pro Ala Thr Ala Thr Pro Thr Pro Thr Ala Thr Ala Ala Ala Ala 255 Val Ala Thr Asn Ala Ala Pro Val Ala Ala Gly Gly Tyr Lys Ile 270 265 <210> 59 <211> 20 <212> PRT <213> Escherichia coli <220> <223> all occurrences of Xaa=hydroxyproline <400> 59 Ala Asp Ala Gly Tyr Thr Xaa Ala Ala Ala Ala Thr Xaa Ala Thr Xaa 5 Ala Ala Thr Xaa 20

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<223> all occurrences of Xaa=hydroxyproline

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Ala Thr Thr Asp 20